

03/20/2002 #10



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RAW SEQUENCE LISTING

DATE: 03/26/2002

PATENT APPLICATION: US/09/811,367B

TIME: 16:17:46

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF3\03262002\I811367B.raw

3 <110> APPLICANT: GEMINI SCIENCE, INC.
 4 Takahashi, Nobuaki
 5 Mikayama, Toshifumi
 7 <120> TITLE OF INVENTION: SOLUBLE MAST CELL FUNCTION ASSOCIATED ANTIGEN (MAFA)
 8 PHARMACEUTICAL COMPOSITIONS AND METHODS OF MAKING AND USING THEM
 10 <130> FILE REFERENCE: 021286/0278719
 12 <140> CURRENT APPLICATION NUMBER: 09/811,367B
 C--> 13 <141> CURRENT FILING DATE: 2002-03-12
 15 <150> PRIOR APPLICATION NUMBER: 60/190,716
 16 <151> PRIOR FILING DATE: 2000-03-17
 18 <160> NUMBER OF SEQ ID NOS: 20
 20 <170> SOFTWARE: PatentIn version 3.0
 22 <210> SEQ ID NO: 1
 23 <211> LENGTH: 189
 24 <212> TYPE: PRT
 25 <213> ORGANISM: Homo sapiens
 27 <400> SEQUENCE: 1
 29 Met Thr Asp Ser Val Ile Tyr Ser Met Leu Glu Leu Pro Thr Ala Thr
 30 1 5 10 15
 32 Gln Ala Gln Asn Asp Tyr Gly Pro Gln Gln Lys Ser Ser Ser Ser Lys
 33 20 25 30
 35 Pro Ser Cys Ser Cys Leu Val Ala Ile Thr Leu Gly Leu Leu Thr Ala
 36 35 40 45
 38 Val Leu Leu Ser Val Leu Leu Tyr Gln Trp Ile Leu Cys Gln Gly Ser
 39 50 55 60
 41 Asn Tyr Ser Thr Cys Ala Ser Cys Pro Ser Cys Pro Asp Arg Trp Met
 42 65 70 75 80
 44 Lys Tyr Gly Asn His Cys Tyr Tyr Phe Ser Val Glu Glu Lys Asp Trp
 45 85 90 95
 47 Asn Ser Ser Leu Glu Phe Cys Leu Ala Arg Asp Ser His Leu Leu Val
 48 100 105 110
 50 Ile Thr Asp Asn Gln Glu Met Ser Leu Leu Gln Val Phe Leu Ser Glu
 51 115 120 125
 53 Ala Phe Cys Trp Ile Gly Leu Arg Asn Asn Ser Gly Trp Arg Trp Glu
 54 130 135 140
 56 Asp Gly Ser Pro Leu Asn Phe Ser Arg Ile Ser Ser Asn Ser Phe Val
 57 145 150 155 160
 59 Gln Thr Cys Gly Ala Ile Asn Lys Asn Gly Leu Gln Ala Ser Ser Cys
 60 165 170 175
 62 Glu Val Pro Leu His Gly Val Cys Lys Lys Val Arg Leu
 63 180 185
 66 <210> SEQ ID NO: 2
 67 <211> LENGTH: 570

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68 <212> TYPE: DNA
69 <213> ORGANISM: Homo sapiens
71 <400> SEQUENCE: 2
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75 gactacggac cacagcaaaa atcttcctct tccaagcctt cttgttcttg ccttgtggca      120
77 ataactttgg ggcttctgac tgcagttctt ctgagtgtgc tgctatacca gtggatcctg      180
79 tgccagggct ccaactactc cacttgtgcc agctgtccta gctgccaga ccgctggatg      240
81 aaatatggta accattgtta ttatttctca gtggaggaaa aggactggaa ttctagtctg      300
83 gaattctgcc tagccagaga ctcacacctc cttgtgataa cggacaatca ggaaatgagc      360
85 ctgctccaag ttttctcag tgaggccttt tgctggattg gtctgaggaa caattctggc      420
87 tggaggtggg aagacggatc acctctaaac ttctcaagga tttcttctaa tagctttgtg      480
89 cagacatgcg gtgccatcaa caaaaatggt cttcaagcct caagctgtga agttccttta      540
91 cacgggtgtg gtaagaaggt cagactttga                               570
95 <210> SEQ ID NO: 3
96 <211> LENGTH: 188
97 <212> TYPE: PRT
98 <213> ORGANISM: Mus musculus
100 <400> SEQUENCE: 3
102 Met Ala Asp Ser Ser Ile Tyr Ser Thr Leu Glu Leu Pro Glu Ala Pro
103 1 5 10 15
105 Gln Val Gln Asp Glu Ser Arg Trp Lys Leu Lys Ala Val Leu His Arg
106 20 25 30
108 Pro His Leu Ser Arg Phe Ala Met Val Ala Leu Gly Leu Leu Thr Val
109 35 40 45
111 Ile Leu Met Ser Leu Leu Met Tyr Gln Arg Ile Leu Cys Cys Gly Ser
112 50 55 60
114 Lys Asp Ser Thr Cys Ser His Cys Pro Ser Cys Pro Ile Leu Trp Thr
115 65 70 75 80
117 Arg Asn Gly Ser His Cys Tyr Tyr Phe Ser Met Glu Lys Lys Asp Trp
118 85 90 95
120 Asn Ser Ser Leu Lys Phe Cys Ala Asp Lys Gly Ser His Leu Leu Thr
121 100 105 110
123 Phe Pro Asp Asn Gln Gly Val Lys Leu Phe Gly Glu Tyr Leu Gly Gln
124 115 120 125
126 Asp Phe Tyr Trp Ile Gly Leu Arg Asn Ile Asp Gly Trp Arg Trp Glu
127 130 135 140
129 Gly Gly Pro Ala Leu Ser Leu Arg Ile Leu Thr Asn Ser Leu Ile Gln
130 145 150 155 160
132 Arg Cys Gly Ala Ile His Arg Asn Gly Leu Gln Ala Ser Ser Cys Glu
133 165 170 175
135 Val Ala Leu Gln Trp Ile Cys Lys Lys Val Leu Tyr
136 180 185
139 <210> SEQ ID NO: 4
140 <211> LENGTH: 997
141 <212> TYPE: DNA
142 <213> ORGANISM: Mus musculus
144 <400> SEQUENCE: 4
146 gtccctcatg gtgtttctca cccacttac agcccacatt cccactgag tgtgaaaggg      60
148 atttggtaga gatggctgac agctctatct attcaacact agagctgccg gaggcacctc      120

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150 aagtccaaga tgagtccaga tggaagctca aagctgtctt acaccggccc catctttccc 180
152 gctttgcaat ggtggctttg gggcttttga ctgtgattct catgagtcta ctgatgtatc 240
154 aacggatcct gtgctgcggc tccaaggact ctacatgttc ccaactgccc agctgcccc 300
156 tcctctggac gaggaatggt agccactgtt actatttttc aatggagaaa aaggactgga 360
158 attctagtct gaaattctgt gcagacaaag gctcacatct ccttacattt ccggacaacc 420
160 agggagtga gctgtttgga gaggacctgg gtcaggactt ttactggatc ggcttgagga 480
162 acattgatgg ctggagggtg gaaggcggcc cagcgctcag cttgaggatt cttaccaaca 540
164 gcttgataca gaggtgcggt gccattcaca gaaatggcct ccaagcctcc agttgtgaag 600
166 ttgctttgca gtggatctgt aagaagggtcc tatactgatg gatgccactg tgtcctgagc 660
168 ctcggaatct ccacatgtgt ttaaaaagag ggaatgggtc tggggaatct ttgtctacaa 720
170 atgtgtgttt aacaaatgcc aaacctgtta tgatatgcca ttagacagag gattagcata 780
172 cctttctggg ggttggcctt ttctgttggt gctttttcgc cgactgttta agtattaggc 840
174 tagccattta aagcctaaat ctgggcaaat caaatgataa agcttattta atggataccc 900
176 accctgcaga tagccaccct ggctctctca tcttctctct gccatctctg tcaagagaga 960
178 gaaactatca tcttcagaga tgaccctgcg catcaga 997

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181 <210> SEQ ID NO: 5

182 <211> LENGTH: 188

183 <212> TYPE: PRT

184 <213> ORGANISM: Rattus norvegicus

186 <400> SEQUENCE: 5

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188 Met Ala Asp Asn Ser Ile Tyr Ser Thr Leu Glu Leu Pro Ala Ala Pro
189 1 5 10 15
191 Arg Val Gln Asp Asp Ser Arg Trp Lys Val Lys Ala Val Leu His Arg
192 20 25 30
194 Pro Cys Val Ser Tyr Leu Val Met Val Ala Leu Gly Leu Leu Thr Val
195 35 40 45
197 Ile Leu Met Ser Leu Leu Leu Tyr Gln Arg Thr Leu Cys Cys Gly Ser
198 50 55 60
200 Lys Gly Phe Met Cys Ser Gln Cys Ser Arg Cys Pro Asn Leu Trp Met
201 65 70 75 80
203 Arg Asn Gly Ser His Cys Tyr Tyr Phe Ser Met Glu Lys Arg Asp Trp
204 85 90 95
206 Asn Ser Ser Leu Lys Phe Cys Ala Asp Lys Gly Ser His Leu Leu Thr
207 100 105 110
209 Phe Pro Asp Asn Gln Gly Val Asn Leu Phe Gln Glu Tyr Val Gly Glu
210 115 120 125
212 Asp Phe Tyr Trp Ile Gly Leu Arg Asp Ile Asp Gly Trp Arg Trp Glu
213 130 135 140
215 Asp Gly Pro Ala Leu Ser Leu Ser Ile Leu Ser Asn Ser Val Val Gln
216 145 150 155 160
218 Lys Cys Gly Thr Ile His Arg Cys Gly Leu His Ala Ser Ser Cys Glu
219 165 170 175
221 Val Ala Leu Gln Trp Ile Cys Glu Lys Val Leu Pro
222 180 185

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224 <210> SEQ ID NO: 6

225 <211> LENGTH: 1461

226 <212> TYPE: DNA

227 <213> ORGANISM: Rattus norvegicus

229 <400> SEQUENCE: 6

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231 caccctgctt actgtcgtca ctccctgctg agtgtgaagg gcgttgggtg gagatggccg      60
233 acaactctat ctactcaaca ttagagctgc ctgctgcacc tcgagtccaa gatgactcca      120
235 gatggaaggt caaagctgtc ttacacgcac cctgtgtttc ctaccttggt atggtggctt      180
237 tggggctttt gactgtgatt ctcatgagtc tactgttgta ccaacggact ctgtgctgtg      240
239 gctccaaggg ctttatgtgt tcccagtgct ccaggtgccc caacctctgg atgaggaacg      300
241 ggagccactg ttactacttc tcaatggaga aaagggactg gaactctagt ctgaagtctt      360
243 gtgcagacaa aggtctgcac ctccctacac ttccggacaa ccaggggagt aacctgttcc      420
245 aggagtatgt gggcgaggac ttttactgga ttggcttgag ggacatcgat ggctggagggt      480
247 gggaagatgg cccagctctc agcttaagca ttctctctaa cagcgtggta cagaagtgtg      540
249 gcaccatcca cagggtgtgg ctccacgcct ccagttgtga ggttgctttg cagtggatct      600
251 gtgagaaggt cctgccctga aggattccac tgtgtcccaa gcctcagatc tgccacatgt      660
253 cttcaaaaag agggaatggg catggggaac ctctgttcac aaagggtgtc ttagcaaatg      720
255 ccaaacctgt tatgatatgc cattagacag gcgttagcat tccttccctg gagctggcat      780
257 ttttcaactg ggctttctca gtcattgttag ccatttaaag cctaaatctg ggcaaatgaa      840
259 atagataaaa tttattttga tggtctttac tgcacaaact caccctggct ttctcatccc      900
261 atactctgcc atatctatca aagatatgtg caaaactatt catctgcaga agaacccccca      960
263 ccacggtcaa taacacatta catagacatc gaatagagac agaaaagcaa acacctcctg     1020
265 ttctcactcc tgcttggaag ctgaagtagc tcaagcctga ggtgtaggga gaagtgcagt     1080
267 ggttaccaga gtccaggaga ctgaaggagt ggtagagggt ggttaatggt ttggctgggt     1140
269 tggggtgacc atcatgatta atgattgttg tatgtttgcc aatatgttgt gaacttccgg     1200
271 atagcgaggt ggaaggaccg tgggtgttac caaatgcctg caggagagat gtgctgagaa     1260
273 cctcgactgg atgatttcca cacacattga aatatcacac tgtgccccat aaatgtgtac     1320
275 aatcattatc tatccctaata ttccctaaaa attaaagaag tcccaattaa aataaaaaaat     1380
277 acctttctgc taaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa     1440
279 aaaaaaaaaa aaaaaaaaaa a      1461
283 <210> SEQ ID NO: 7
284 <211> LENGTH: 32
285 <212> TYPE: DNA
286 <213> ORGANISM: Artificial Sequence
288 <220> FEATURE:
289 <223> OTHER INFORMATION: Description of Artificial Sequence:Primer
291 <400> SEQUENCE: 7
293 ccttgtgatg gtggctttgg ggcttttgac tg      32
296 <210> SEQ ID NO: 8
297 <211> LENGTH: 28
298 <212> TYPE: DNA
299 <213> ORGANISM: Artificial Sequence
301 <220> FEATURE:
302 <223> OTHER INFORMATION: Description of Artificial Sequence:Primer
304 <400> SEQUENCE: 8
306 actgcaaagc aacctcacia ctggaggc      28
309 <210> SEQ ID NO: 9
310 <211> LENGTH: 30
311 <212> TYPE: DNA
312 <213> ORGANISM: Artificial Sequence
314 <220> FEATURE:
315 <223> OTHER INFORMATION: Description of Artificial Sequence:Primer
317 <400> SEQUENCE: 9
319 atatgatcc tccaaggact ctacatgttc      30

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322 <210> SEQ ID NO: 10
323 <211> LENGTH: 34
324 <212> TYPE: DNA
325 <213> ORGANISM: Artificial Sequence
327 <220> FEATURE:
328 <223> OTHER INFORMATION: Description of Artificial Sequence:Primer
330 <400> SEQUENCE: 10
332 atatgcggcc gctcagtata ggacotttctt acag          34
335 <210> SEQ ID NO: 11
336 <211> LENGTH: 64
337 <212> TYPE: DNA
338 <213> ORGANISM: Artificial Sequence
340 <220> FEATURE:
341 <223> OTHER INFORMATION: Description of Artificial Sequence:Primer
343 <400> SEQUENCE: 11
345 cccggatccg catcaccatc accatcacgc ggccgcttcc aaggactota catgttccca    60
347 ctgc                                                                    64
350 <210> SEQ ID NO: 12
351 <211> LENGTH: 34
352 <212> TYPE: DNA
353 <213> ORGANISM: Artificial Sequence
355 <220> FEATURE:
356 <223> OTHER INFORMATION: Description of Artificial Sequence:Primer
358 <400> SEQUENCE: 12
360 atatgcggcc gctcagtata ggacotttctt acag          34
363 <210> SEQ ID NO: 13
364 <211> LENGTH: 73
365 <212> TYPE: DNA
366 <213> ORGANISM: Artificial Sequence
368 <220> FEATURE:
369 <223> OTHER INFORMATION: Description of Artificial Sequence:Primer
371 <400> SEQUENCE: 13
373 cccaagctta caaccatggc tgaccgctct atcgccctcaa cagccgagct gccggaggca    60
375 cctcaagtcc aag                                                                    73
378 <210> SEQ ID NO: 14
379 <211> LENGTH: 66
380 <212> TYPE: DNA
381 <213> ORGANISM: Artificial Sequence
383 <220> FEATURE:
384 <223> OTHER INFORMATION: Description of Artificial Sequence:Primer
386 <400> SEQUENCE: 14
388 cccctcgagc tacagatcct cttcagagat gagttttctgc tcgtatagga ctttcttaca    60
390 gatcca                                                                    66
393 <210> SEQ ID NO: 15
394 <211> LENGTH: 32
395 <212> TYPE: DNA
396 <213> ORGANISM: Artificial Sequence
398 <220> FEATURE:
399 <223> OTHER INFORMATION: Description of Artificial Sequence:Primer

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VERIFICATION SUMMARY

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L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date